

C. Kaufman

#14

1646

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/954,771

DATE: 10/08/1999
TIME: 14:55:28

INPUT SET: S33602.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages

ENTERED

SEQUENCE LISTING

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Ingham, Phillip W.
6 McMahon, Andrew P.
7 Tabin, Clifford J.
8
9 (ii) TITLE OF INVENTION: Vertebrate Embryonic Pattern-Inducing
10 Proteins and Uses Related Thereto
11
12 (iii) NUMBER OF SEQUENCES: 48
13
14 (iv) CORRESPONDENCE ADDRESS:
15 (A) ADDRESSEE: FOLEY, HOAG & ELIOT LLP
16 (B) STREET: One Post Office Square
17 (C) CITY: Boston
18 (D) STATE: MA
19 (E) COUNTRY: USA
20 (F) ZIP: 02109-2170
21
22 (v) COMPUTER READABLE FORM:
23 (A) MEDIUM TYPE: Floppy disk
24 (B) COMPUTER: IBM PC compatible
25 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
26 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
27
28 (vi) CURRENT APPLICATION DATA:
29 (A) APPLICATION NUMBER: US 08/954,771
30 (B) FILING DATE: 20-OCT-1997
31
32 (vii) PRIOR APPLICATION DATA:
33 (A) APPLICATION NUMBER: US 08/462,386
34 (B) FILING DATE: 05-JUN-1995
35
36 (vii) PRIOR APPLICATION DATA:
37 (A) APPLICATION NUMBER: US 08/435,093
38 (B) FILING DATE: 04-MAY-1995
39
40 (vii) PRIOR APPLICATION DATA:
41 (A) APPLICATION NUMBER: US 08/356,060
42 (B) FILING DATE: 14-DEC-1994
43
44 (vii) PRIOR APPLICATION DATA:
45 (A) APPLICATION NUMBER: US 08/176,427
46 (B) FILING DATE: 30-DEC-1993

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/954,771DATE: 10/08/1999
TIME: 14:55:28

INPUT SET: S33602.raw

47
48 (viii) ATTORNEY/AGENT INFORMATION:
49 (A) NAME: Vincent, Matthew P.
50 (B) REGISTRATION NUMBER: 36,709
51 (C) REFERENCE/DOCKET NUMBER: HMV-006.11
52
53 (ix) TELECOMMUNICATION INFORMATION:
54 (A) TELEPHONE: 617-832-1000
55 (B) TELEFAX: 617-832-7000
56
57
58 (2) INFORMATION FOR SEQ ID NO:1:
59
60 (i) SEQUENCE CHARACTERISTICS:
61 (A) LENGTH: 1277 base pairs
62 (B) TYPE: nucleic acid
63 (C) STRANDEDNESS: both
64 (D) TOPOLOGY: linear
65
66 (ii) MOLECULE TYPE: cDNA
67
68
69 (ix) FEATURE:
70 (A) NAME/KEY: CDS
71 (B) LOCATION: 1..1275
72
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74 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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97 Pro Asn Tyr Asn Pro Asp Ile Ile Phe Lys Asp Glu Glu Asn Thr Gly
98 85 90 95
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/954,771

DATE: 10/08/1999
TIME: 14:55:28

INPUT SET: S33602.raw

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105	Ala Ile Ser Val Met Asn Gln Trp Pro Gly Val Lys Leu Arg Val Thr	
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109	Glu Gly Trp Asp Glu Asp Gly His His Ser Glu Glu Ser Leu His Tyr	
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RAW SEQUENCE LISTING PATENT APPLICATION US/08/954,771

DATE: 10/08/1999
TIME: 14:55:29

INPUT SET: S33602.raw

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161 Leu Thr Ala Gln Gly Thr Ile Leu Ile Asn Arg Val Leu Ala Ser Cys
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165 Tyr Ala Val Ile Glu Glu His Ser Trp Ala His Trp Ala Phe Ala Pro
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168 TTC CGC TTG GCT CAG GGG CTG CTG GCC GCC CTC TGC CCA GAT GGG GCC 1152
169 Phe Arg Leu Ala Gln Gly Leu Leu Ala Ala Leu Cys Pro Asp Gly Ala
170 370 375 380
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173 Ile Pro Thr Ala Ala Thr Thr Thr Thr Gly Ile His Trp Tyr Ser Arg
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176 CTC CTC TAC CGC ATC GGC AGC TGG GTG CTG GAT GGT GAC GCG CTG CAT 1248
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185 (2) INFORMATION FOR SEQ ID NO:2:
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187 (i) SEQUENCE CHARACTERISTICS:
188 (A) LENGTH: 1190 base pairs
189 (B) TYPE: nucleic acid
190 (C) STRANDEDNESS: both
191 (D) TOPOLOGY: linear
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193 (ii) MOLECULE TYPE: cDNA
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196 (ix) FEATURE:
197 (A) NAME/KEY: CDS
198 (B) LOCATION: 1..1188
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201 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
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212	Arg	Arg	Tyr	Val	Arg	Lys	Gln	Leu	Val	Pro	Leu	Leu	Tyr	Lys	Gln	Phe		
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216	Val	Pro	Ser	Met	Pro	Glu	Arg	Thr	Leu	Gly	Ala	Ser	Gly	Pro	Ala	Glu		
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227	CGC	CTG	ATG	ACA	GAG	CGT	TGC	AAA	GAG	CGG	GTG	AAC	GCT	CTA	GCC	ATC		336
228	Arg	Leu	Met	Thr	Glu	Arg	Cys	Lys	Glu	Arg	Val	Asn	Ala	Leu	Ala	Ile		
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244	Leu	Leu	Ala	Arg	Leu	Ala												

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/954,771

DATE: 10/08/1999
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INPUT SET: S33602.raw

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